

REPLACEMENT SHEET

Fig. 15A

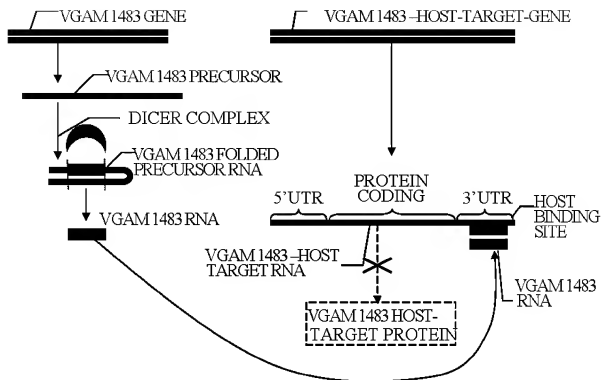


Fig. 15B

GTGTGACGTGAGTCTGATCCAACACTGAACGCTTTCGTC SEQ ID: 37404
 GTGTTTTTCATGCAGCTTTTACAGACCATGACAAGCCTGA
CGAGAGCGTTCATCGGGGCATGAAGTACGCATTAC

TGACAAGCCTGACGAGAGCGTTCA

SEQ ID: 37405

Fig. 15C

--	AG	-	A	AACAC		T	T	CAG	TT
GTGTGA	CGTG	TC	TG	TCC		TGAACGCTTTCGTCG	GTTT	TCATG	CT T
CACATT	GCAT	AG	AC	GGG		ACTTGCAGAGCAGT	CGAA	AGTAC	GA /
AC	GA	T	-	GCT--		<u>C</u>	<u>C</u>	CA-	CA

REPLACEMENT SHEET

Fig. 15D/1

		CAA	CGA		
	5'	TGA	GCCTGA	GAGCGTTCA	3'
C7					
BINDING SITE	3'	ACT	CGGACT	CTCGTAAGT	5'
		ATC	TAC		
		A	ACGA	G	
	5'	TG	CAAGCCTG	GA CGTTCA	3'
DUSP5					
BINDING SITE	3'	AC	GTTCGGAC	CT GTGAGT	5'
		-	CA--	G	
		A	AC		
	5'	TG	CAAGCCTG	GAGAG	3'
ATP6V1F					
BINDING SITE	3'	AC	GTTCGGAC	CTCTC	5'
		C	CC		
		CAA	AGAGCG		
	5'	TGA	GCCTGACG	TTCA	3'
DKFZP564C103					
BINDING SITE	3'	ACT	CGGACTGC	AAGT	5'
		AC-	GAGAG-		
		CA	C	C	
	5'	GA	AGC TGACGAGAG	GT	3'
KIAA0561					
BINDING SITE	3'	CT	TCG ACTGCTCTC	CA	5'
		AC	A	C	
		A	A G A		
	5'	TG	CAAGCCTG C AG GC		3'
KIAA0935					
BINDING SITE	3'	AC	GTTCGGAC G TC CG		5'
		C	C - C		
			AGC		
	5'	GACAAGCCTGACGAG	GTTCA		3'
MIC2L1					
BINDING SITE	3'	CTGTTCGGACTGTTT	CAAGT		5'

		C	C-		
	5'	TGACAAG CTGA	GAGAGCG		3'
PPP4R1L					
BINDING SITE	3'	ACTGTTC GACT	TTCTCGC		5'
		A	AC		
			TGACG	GC-	
	5'	TGACAAGCC	AGA GTTC		3'
ZAK					
BINDING SITE	3'	ACTGTTCGG	TCT CAAG		5'
		-----	AGT		

REPLACEMENT SHEET

Fig. 15D/2

		CC ACGA		
LOC149566	5'	TGACAAG TG	GAGCGTT	3' SEQ ID:37405
BINDING SITE	3'	ACTGTTC AC	CTTGCAA	5' SEQ ID:37415
		A- CCA-		
		A AC G CG		
LOC150368	5'	TG CAAGCCTG GA AG	TTC	3' SEQ ID:37405
BINDING SITE	3'	AC GTTCGGAC CT TC	GAG	5' SEQ ID:37416
		C A- G A-		
		A CGAGA C		
LOC255718	5'	TGACA GCCTGA	G GTTC	3' SEQ ID:37405
BINDING SITE	3'	ACTGT CGGACT	C CAGG	5' SEQ ID:37417
		C ----- C		